

## Quick guide

# Presenilins

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**What are they?** Presenilins (PS) have eight transmembrane domains, and a cytosolic loop which undergoes self-cleavage to generate amino- and carboxy-terminal fragments that remain associated in a high molecular weight complex. There are two *PS* genes in mammals and homologs in *Caenorhabditis elegans*, *Drosophila melanogaster*, and *Arabidopsis thaliana* but not in *Saccharomyces cerevisiae*.

**Where are they found?** They are localized in the endoplasmic reticulum and Golgi membranes, and have been detected at the plasma membrane. They are expressed in most cell types throughout development and in adulthood.

**How did they get that name?** Presenilins were identified as susceptibility genes for familial Alzheimer's disease.

**What do they do normally?** Analyses of the *C. elegans* orthologs Sel-12 and Hop-1, and of double knockout mice showed that presenilins are involved in Notch signalling and particularly in the cleavage of the trans-membrane domain of the Notch receptor after ligand binding, releasing an intracellular fragment that allows signal transduction. This activity is called  $\gamma$ -secretase by analogy with cleavage of the amyloid precursor protein, APP, as presenilins are involved in the proteolysis of APP. In this case,  $\beta$ -secretase produces a membrane-associated carboxy-terminal fragment of APP, which is cleaved by  $\gamma$ -secretase. The  $\gamma$ -secretase cleavage of both APP and Notch is facilitated by PS1 and PS2, and there is evidence that the presenilins themselves could be the proteases that mediate this event.

**And in Alzheimer's disease?** Amyloid  $\beta$  peptides 40/42 are normal products of APP metabolism. The appearance of Alzheimer's disease correlates with increased APP cleavage, and a change in the ratio of the two products which may lead to the formation of so-called senile plaques.

### Other putative roles for presenilins...

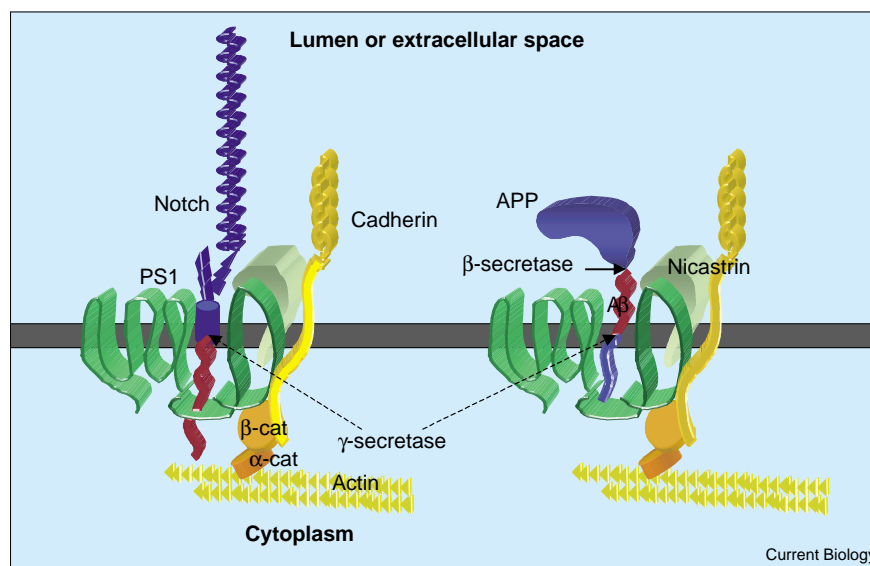
They may interact with cadherin-catenin complexes to modulate the canonical Wnt pathway, and participate in cell-cell contacts. They may also interact with calsenilin to modulate cellular calcium metabolism and control neuronal apoptosis. Presenilins may be required for a normal Unfolded Protein Response and may act as general regulators of apoptosis and as chaperones.

**Can we do without them?** No, although PS1 and PS2 are partially redundant. *PS1*<sup>-/-</sup> mice die shortly after birth with defects of the axial skeleton, neurogenesis and neuronal survival while *PS2*<sup>-/-</sup> mice show very mild defects. Mice lacking both genes have a phenotype similar to full Notch-1 deficiency, and die during early embryogenesis. On the other hand, too much presenilin is also detrimental, as in Alzheimer's disease.

### Where can I find out more?

<http://www.alzforum.org>  
Steiner H, Haas C: Intramembrane proteolysis by presenilins. *Nat Rev* 2000, 1:217-224.  
Kopan R, Goate A: A common enzyme connects Notch signaling and Alzheimer's disease. *Genes Dev* 2000, 14:2799-2806.

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A model for presenilin-containing  $\gamma$ -secretase complex. Heterodimeric presenilin (green) interacts directly with the  $\gamma$ -secretase substrates Notch (left part) or APP (right part), and also with nicastrin (light green), cadherin (yellow) and  $\beta$ -catenin ( $\beta$ -cat,

orange). The yellow- and orange-colored items represent proteins that are linked to the cytoskeleton. In red are the intracytoplasmic part of the Notch receptor and the amyloid  $\beta$  ( $A\beta$ ) peptides derived from APP, as a result of  $\gamma$ -secretase activity.